1601

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TECH CENTER 1600/2900

RAW SEQUENCE LISTING DATE: 12/18/2001 PATENT APPLICATION: US/09/783,338A TIME: 16:37:40

Input Set : N:\Crf3\RULE60\09783338A.raw
Output Set: N:\CRF3\12182001\1783338A.raw

SEQUENCE LISTING

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(1) GENERAL INFORMATION:
      5
              (i) APPLICANT: Glazer, Peter M.
                             Havre, Pamela A.
      6
      8
            (ii) TITLE OF INVENTION: Chemically Modified Oligonucleotide for
      9
                                      Site-Directed Mutagenesis
     11
           (iii) NUMBER OF SEQUENCES: 13
     13
            (iv) CORRESPONDENCE ADDRESS:
     14
                   (A) ADDRESSEE: Patrea L. Pabst
     15
                   (B) STREET: 1100 Peachtree Street, Suite 2800
                   (C) CITY: Atlanta
     16
                   (D) STATE: Georgia
     17
                                                                ENTERED
     18
                   (E) COUNTRY: USA
     19
                   (F) ZIP: 30309-4530
     21
             (V) COMPUTER READABLE FORM:
     22
                   (A) MEDIUM TYPE: Floppy disk
     23
                   (B) COMPUTER: IBM PC compatible
     24
                   (C) OPERATING SYSTEM: PC-DOS/MS-DOS
     25
                   (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
     27
            (vi) CURRENT APPLICATION DATA:
C--> 28
                   (A) APPLICATION NUMBER: US/09/783,338A
C--> 29
                   (B) FILING DATE: 14-Feb-2001
     30
                   (C) CLASSIFICATION:
     32
           (vii) PRIOR APPLICATION DATA:
     33
                   (A) APPLICATION NUMBER: 08/083,088
     34
                   (B) FILING DATE: 25-JUN-1993
     37
          (viii) ATTORNEY/AGENT INFORMATION:
     38
                   (A) NAME: Pabst, Patrea L.
     39
                   (B) REGISTRATION NUMBER: 31,284
     40
                   (C) REFERENCE/DOCKET NUMBER: YU109
     43
            (ix) TELECOMMUNICATION INFORMATION:
     44
                   (A) TELEPHONE: (404)-815-6508
     45
                  (B) TELEFAX: (404)-815-6555
        (2) INFORMATION FOR SEQ ID NO: 1:
     47
     49
             (i) SEQUENCE CHARACTERISTICS:
     50
                   (A) LENGTH: 11 base pairs
     51
                  (B) TYPE: nucleic acid
     52
                  (C) STRANDEDNESS: single
     53
                  (D) TOPOLOGY: linear
     55
            (ii) MOLECULE TYPE: DNA (genomic)
     57
           (iii) HYPOTHETICAL: NO
     59
            (iv) ANTI-SENSE: NO
     61
            (ix) FEATURE:
     62
                  (A) NAME/KEY: misc_feature
     63
                  (B) LOCATION: 1..11
                  (D) OTHER INFORMATION: /function= "N is
     65 4'-hydroxymethyl-4,5',8-trimethylpsoralen"
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Input Set : N:\Crf3\RULE60\09783338A.raw
Output Set: N:\CRF3\12182001\1783338A.raw

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JAN 0 8 2002
67
         (x) PUBLICATION INFORMATION:
              (A) AUTHORS: Glazer, Peter M.
                                                                         TECH CENTER 1600/2900
69 Havre, Pamela A.
70
              (K) RELEVANT RESIDUES IN SEQ ID NO:1: FROM 1 TO 11
72
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
74 NAGGAAGGGG G
                                                                              11
80 (2) INFORMATION FOR SEQ ID NO: 2:
82
        (i) SEQUENCE CHARACTERISTICS:
83
              (A) LENGTH: 10 base pairs
84
              (B) TYPE: nucleic acid
85
              (C) STRANDEDNESS: single
86
              (D) TOPOLOGY: linear
88
       (ii) MOLECULE TYPE: DNA (genomic)
90
      (iii) HYPOTHETICAL: NO
92
       (iv) ANTI-SENSE: NO
94
       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:
96 AGGAAGGGGG
                                                                              10
98 (2) INFORMATION FOR SEQ ID NO: 3:
100
         (i) SEQUENCE CHARACTERISTICS:
101
               (A) LENGTH: 10 base pairs
102
               (B) TYPE: nucleic acid
103
               (C) STRANDEDNESS: single
104
               (D) TOPOLOGY: linear
106
        (ii) MOLECULE TYPE: DNA (genomic)
108
       (iii) HYPOTHETICAL: NO
110
        (iv) ANTI-SENSE: NO
112
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:
114 GGGGGAAGGA
                                                                               10
117 (2) INFORMATION FOR SEO ID NO: 4:
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              (A) LENGTH: 8 base pairs
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              (C) STRANDEDNESS: single
123
              (D) TOPOLOGY: linear
125
        (ii) MOLECULE TYPE: DNA (genomic)
127
       (iii) HYPOTHETICAL: NO
129
        (iv) ANTI-SENSE: NO
131
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133 CCCCCTTC
                                                                                8
135 (2) INFORMATION FOR SEQ ID NO: 5:
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              (B) TYPE: nucleic acid
140
              (C) STRANDEDNESS: single
141
              (D) TOPOLOGY: linear
143
        (ii) MOLECULE TYPE: DNA (genomic)
145
       (iii) HYPOTHETICAL: NO
147
        (iv) ANTI-SENSE: NO
149
        (ix) FEATURE:
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Input Set : N:\Crf3\RULE60\09783338A.raw
Output Set: N:\CRF3\12182001\1783338A.raw

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150
               (A) NAME/KEY: misc_feature
151
               (B) LOCATION: 1..11
152
               (D) OTHER INFORMATION: /function= "N is
153 4'-hydroxymethyl-4,5',8-trimethylpsoralen"
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:
157 NGGGGGAAGG A
                                                                               11
159 (2) INFORMATION FOR SEQ ID NO: 6:
         (i) SEQUENCE CHARACTERISTICS:
162
               (A) LENGTH: 18 base pairs
               (B) TYPE: nucleic acid
163
164
               (C) STRANDEDNESS: single
165
               (D) TOPOLOGY: linear
167
        (ii) MOLECULE TYPE: DNA (genomic)
       (iii) HYPOTHETICAL: NO
169
171
        (iv) ANTI-SENSE: NO
173
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:
175 GGTTCGAATC CTTCCCCC
                                                                              18
177 (2) INFORMATION FOR SEQ ID NO: 7:
179
         (i) SEQUENCE CHARACTERISTICS:
180
              (A) LENGTH: 18 base pairs
              (B) TYPE: nucleic acid
181
182
              (C) STRANDEDNESS: single
183
              (D) TOPOLOGY: linear
185
        (ii) MOLECULE TYPE: DNA (genomic)
187
       (iii) HYPOTHETICAL: NO
189
        (iv) ANTI-SENSE: NO
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:
195 GGTTCGAAAC CTTCCCCC
                                                                              18
197 (2) INFORMATION FOR SEQ ID NO: 8:
199
         (i) SEQUENCE CHARACTERISTICS:
200
              (A) LENGTH: 35 base pairs
201
              (B) TYPE: nucleic acid
202
              (C) STRANDEDNESS: single
203
              (D) TOPOLOGY: linear
205
        (ii) MOLECULE TYPE: DNA (genomic)
207
       (iii) HYPOTHETICAL: NO
209
        (iv) ANTI-SENSE: NO
211
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:
213 TGGTGGTGGG GGAAGGATTC GAACCTTCGA AGTCG
                                                                              35
215 (2) INFORMATION FOR SEQ ID NO: 9:
217
         (i) SEQUENCE CHARACTERISTICS:
218
              (A) LENGTH: 35 base pairs
219
              (B) TYPE: nucleic acid
220
              (C) STRANDEDNESS: single
221
              (D) TOPOLOGY: linear
223
        (ii) MOLECULE TYPE: DNA (genomic)
225
       (iii) HYPOTHETICAL: NO
227
        (iv) ANTI-SENSE: NO
231
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:
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Input Set : N:\Crf3\RULE60\09783338A.raw
Output Set: N:\CRF3\12182001\1783338A.raw

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233 CGACTTCGAA GGTTCGAATC CTTCCCCCAC CACCA
                                                                               35
235 (2) INFORMATION FOR SEQ ID NO: 10:
237
          (i) SEQUENCE CHARACTERISTICS:
238
               (A) LENGTH: 121 base pairs
239
               (B) TYPE: nucleic acid
240
               (C) STRANDEDNESS: single
241
               (D) TOPOLOGY: linear
243
        (ii) MOLECULE TYPE: DNA (genomic)
245
       (iii) HYPOTHETICAL: NO
247
        (iv) ANTI-SENSE: NO
249
        (ix) FEATURE:
250
               (A) NAME/KEY: misc_feature
251
               (B) LOCATION: 1..60
252
               (D) OTHER INFORMATION: /function= "At positions 16 and 57,
253 N is G or T."
255
        (ix) FEATURE:
256
               (A) NAME/KEY: misc_feature
257
               (B) LOCATION: 1..121
258
               (D) OTHER INFORMATION: /function= "At position 17, N is A,
259 G or T."
        (ix) FEATURE:
261
262
               (A) NAME/KEY: misc_feature
263
               (B) LOCATION: 1..121
264
               (D) OTHER INFORMATION: /function= "At position 18, N is A,
265 C or T."
268
        (ix) FEATURE:
269
              (A) NAME/KEY: misc_feature
270
               (B) LOCATION: 1..121
271
               (D) OTHER INFORMATION: /function= "At position 20, N is G
272 or A."
274
       (ix) FEATURE:
275
              (A) NAME/KEY: misc_feature
276
              (B) LOCATION: 1..121
277
              (D) OTHER INFORMATION: /function= "At position 24, N is C
278 or T."
280
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 10:
282 TGGTGGTGGG GGAAGNNNTN GAANCTTCGA AGTCGATGAC GGCAGATTTA GAGTCTNCTC
                                                                              60
284 CCTTTGGCCG CTCGGGAACC CCACCACAGG TAATGCTTTT ACGGGGCGCA TCATATCAAA
                                                                             120
286 T
                                                                             121
288 (2) INFORMATION FOR SEQ ID NO: 11:
290
         (i) SEQUENCE CHARACTERISTICS:
291
              (A) LENGTH: 121 base pairs
292
              (B) TYPE: nucleic acid
293
              (C) STRANDEDNESS: single
294
              (D) TOPOLOGY: linear
296
        (ii) MOLECULE TYPE: DNA (genomic)
298
       (iii) HYPOTHETICAL: NO
300
        (iv) ANTI-SENSE: NO
302
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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Input Set: N:\Crf3\RULE60\09783338A.raw
Output Set: N:\CRF3\12182001\1783338A.raw

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304 ATTTGATATG ATGCGCCCCG TAAAAGCATT ACCTGTGGTG GGGTTCCCGA GCGGCCAAAG
                                                                               60
 306 GGAGCAGACT CTAAATCTGC CGTCATCGAC TTCGAAGGTT CGAATCCTTC CCCCACCACC
                                                                              120
 308 A
                                                                              121
 310 (2) INFORMATION FOR SEQ ID NO: 12:
 312
          (i) SEQUENCE CHARACTERISTICS:
313
               (A) LENGTH: 121 base pairs
314
               (B) TYPE: nucleic acid
315
               (C) STRANDEDNESS: single
316
               (D) TOPOLOGY: linear
318
         (ii) MOLECULE TYPE: DNA (genomic)
320
        (iii) HYPOTHETICAL: NO
322
         (iv) ANTI-SENSE: NO
325
         (ix) FEATURE:
326
               (A) NAME/KEY: misc_feature
327
               (B) LOCATION: 1..121
328
               (D) OTHER INFORMATION: /function= "At position 7, N is G
329 or T."
331
        (ix) FEATURE:
332
               (A) NAME/KEY: misc_feature
333
              (B) LOCATION: 1..121
334
              (D) OTHER INFORMATION: /function= "At positions 12, 20 and
335 117, N is C or G."
337
        (ix) FEATURE:
               (A) NAME/KEY: misc_feature
338
339
              (B) LOCATION: 1..121
              (D) OTHER INFORMATION: /function= "At positions 15, 16,
341 21, 29, 78, 83 and 108, N is A or G."
342
        (ix) FEATURE:
343
              (A) NAME/KEY: misc_feature
344
              (B) LOCATION: 1..121
345
              (D) OTHER INFORMATION: /function= "At positions 18, 49,
346 68, 69, and 107, N is C or T."
        (ix) FEATURE:
348
              (A) NAME/KEY: misc_feature
349
              (B) LOCATION: 1..121
350
              (D) OTHER INFORMATION: /function= "At position 114, N is
351 C, T or A."
353
        (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 12:
355 TGGTGGNGGG GNAANNANTN NAACCTTCNA AGTCGATGAC GGCAGATTNN GAGTCTNCTC
                                                                              60
357 CCTTTGGNNG CTCGGGANCC CCNCCACAGG TAATGCTTTT ACGGGGNNCA TCANNTNAAA
                                                                             120
                                                                             121
362 (2) INFORMATION FOR SEQ ID NO: 13:
364
         (i) SEQUENCE CHARACTERISTICS:
365
              (A) LENGTH: 121 base pairs
366
              (B) TYPE: nucleic acid
367
              (C) STRANDEDNESS: single
368
              (D) TOPOLOGY: linear
370
        (ii) MOLECULE TYPE: DNA (genomic)
372
       (iii) HYPOTHETICAL: NO
```

VERIFICATION SUMMARY

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DATE: 12/18/2001 TIME: 16:37:41 PATENT APPLICATION: US/09/783,338A

Input Set : N:\Crf3\RULE60\09783338A.raw Output Set: N:\CRF3\12182001\1783338A.raw

L:28 M:220 C: Keyword misspelled or invalid format, [(A) APPLICATION NUMBER:]

L:29 M:220 C: Keyword misspelled or invalid format, [(B) FILING DATE:]